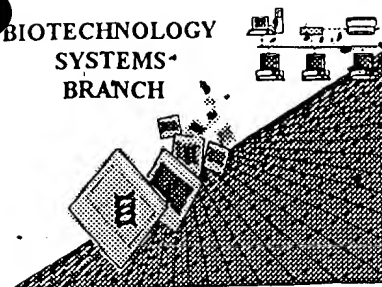


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



0590
1005

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 4-13 09/865579

Source: OIPB

Date Processed by STIC: 11-13-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/865,579

DATE: 11/13/2001
 TIME: 13:48:13

Input Set : A:\es.txt
 Output Set: N:\CRF3\11132001\I865579.raw

3 <110> APPLICANT: TOSOH Corporation
 5 <120> TITLE OF INVENTION: Oligonucleotides and Method for Detection of mecA Gene of Methicillin

6 Resistant Staphylococcus Aureus
 8 <130> FILE REFERENCE: PA211-0187
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/865,579
 C--> 10 <141> CURRENT FILING DATE: 2001-05-29
 E--> 10 <160> NUMBER OF SEQ ID NOS: 17

**Does Not Comply
 Corrected Diskette Needed**

*Field 160 indicates 17
 Found 30 sequences*

ERRORED SEQUENCES

348 <210> SEQ ID NO: 30
 349 <211> LENGTH: 28
 350 <212> TYPE: DNA
 351 <213> ORGANISM: T7 phage
 353 <220> FEATURE:
 354 <221> NAME/KEY: promoter
 355 <223> OTHER INFORMATION: Promoter sequence of T7 polymerase
 357 <400> SEQUENCE: 30
 358 aattctaata cgactcacta tagggaga 28

W--> 359 ??
 W--> 367 - 11 -
 E--> 372 - 1 -

Delete End of file Non ASCII text

Errors

1. Field 160 indicates 17 sequences in the sequence listing.
 There are actually 30 sequences in the sequence listing
2. End of file contains Non ASCII text

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/865,579

DATE: 11/13/2001

TIME: 13:48:14

Input Set : A:\es.txt

Output Set: N:\CRF3\11132001\I865579.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:359 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:367 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:30
L:372 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:28 SEQ:30
L:10 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (17) Counted (30)